
Sequence Listing was accepted with existing errors.

See attached Validation Report.

If you need help call the Patent Electronic Business Center at (866)

217-9197 (toll free).

Reviewer: Durreshwar Anjum

Timestamp: Wed May 23 12:34:02 EDT 2007

Validated By CRFValidator v 1.0.2

Application No: 10584367 Version No: 1.1

Input Set:

Output Set:

Started: 2007-05-23 12:33:50.996

Finished: 2007-05-23 12:33:52.456

Elapsed: 0 hr(s) 0 min(s) 1 sec(s) 460 ms

Total Warnings: 0

Total Errors: 0

No. of SeqIDs Defined: 56

Actual SeqID Count: 56

SEQUENCE LISTING

<110>	Imperial Innovations Limited Tang, Christoph M											
<120>	Identification of Antigenically Important Neisseria Antigens by Screening Insertional Mutant Libraries with Antiserum											
<130>	06-451											
<140>	10/584,367											
<141>	2006-07-26											
	GB04/005441 2004-12-23											
<151>	2004-12-23											
<150>	PCT,	/GB04/005441	1									
<151>												
<160>	56											
<100>	50											
<170>	PatentIn version 3.3											
<210>	1											
<211>	2628											
<212>												
<213>	Neis	sseria menir	ngitidis (g	roup B)								
<400>	1											
	1 rcca	accaactacc	ccgccgatgc	ccgatgatga	cgaaatttac	agactgtacg	60					
9	, 5	999	999	9 9 9	- 9	99						
cggtcaa	acc	gtattcagcc	gccaacccac	aggggataca	tcttgaaaaa	caacagacaa	120					
atcaaac	ctga	ttgccgcctc	cgtcgcagtt	gccgcatcct	ttcaggcaca	tgctggactg	180					
ggcggad	ctga	atatccagtc	caaccttgac	gaaccctttt	ccggcagcat	taccgtaacc	240					
ggcgaaq	gaag	ccaaagccct	gctaggcggc	ggcagcgtta	ccgtttccga	aaaaggcctg	300					
accgcca	aaag	tccacaagtt	gggcgacaaa	gccgtcattg	ccgtttcttc	cgaacaggca	360					
gtaagaq	gatc	ccgtcctggt	gttccgcatc	ggcgcaggcg	cacaggtacg	cgaatacacc	420					
gccatco	ctcg	atcctgtcgg	ctactcgccc	aaaaccaaat	ctgcactttc	agacggcaag	480					
acacaco	cgca	aaaccgctcc	gacagcagag	tcccaagaaa	atcaaaacgc	caaagccctc	540					
cgcaaaa	accg	ataaaaaaga	cagcgcgaac	gcagccgtca	aaccggcata	caacggcaaa	600					
acccata	accg	tccgcaaagg	cgaaacggtc	aaacagattg	ccgccgccat	ccgcccgaaa	660					
cacctga	acgc	tcgaacaggt	tgccgatgcg	ctgctgaagg	caaacccaaa	tgtttccgca	720					
cacggca	agac	tgcgtgcggg	cagcgtgctt	cacattccga	atctgaacag	gatcaaagcg	780					

gaacaaccca aaccgcaaac ggcgaaaccc aaagccgaaa ccgcatccat gccgtccgaa

840

ccgtccaaac	aggcaacggt	agagaaaccg	gttgaaaaac	ctgaagcaaa	agttgccgcg	900
cccgaagcaa	aagcggaaaa	accggccgtt	cgacccgaac	ctgtacccgc	tgcaaatact	960
gccgcatcgg	aaaccgctgc	cgaatccgcc	ccccaagaag	ccgccgcttc	tgccatcgac	1020
acgccgaccg	acgaaaccgg	taacgccgtt	tccgaacctg	tcgaacaggt	ttctgccgaa	1080
gaagaaaccg	aaagcggact	gtttgacggt	ctgttcggcg	gttcgtacac	cttgctgctt	1140
gccggcggag	gcgcggcatt	aatcgccctg	ctgctgcttt	tgegeettge	ccaatccaaa	1200
cgcgcgcgcc	gtaccgaaga	atccgtccct	gaggaagagc	ctgaccttga	cgacgcggca	1260
gacgacggca	tagaaatcac	ctttgccgaa	gtcgaaactc	cggcaacgcc	cgaacccgct	1320
ccgaaaaacg	atgtaaacga	cacacttgcc	ttagatgggg	aatctgaaga	agagttatcg	1380
gcaaaacaaa	cgttcgatgt	cgaaaccgat	acgccttcca	accgcatcga	cttggatttc	1440
gacagcctgg	cagccgcgca	aaacggcatt	ttatccggcg	cacttacgca	ggatgaagaa	1500
acccaaaaac	gcgcggatgc	cgattggaac	gccatcgaat	ccacagacag	cgtgtacgag	1560
cccgagacct	tcaacccgta	caaccctgtc	gaaatcgtca	tcgacacgcc	cgaaccggaa	1620
tetgtegeee	aaactgccga	aaacaaaccg	gaaaccgtcg	ataccgattt	ctccgacaac	1680
ctgccctcaa	acaaccatat	cggcacagaa	gaaacagctt	ccgcaaaacc	tgcctcaccc	1740
tccggactgg	caggetteet	gaaggcttcc	tcgcccgaaa	ccatcttgga	aaaaacagtt	1800
gccgaagtcc	aaacaccgga	agagttgcac	gatttcctga	aagtgtacga	aaccgatgcc	1860
gtcgcggaaa	ctgcgcctga	aacgcccgat	ttcaacgccg	ccgcagacga	tttgtccgca	1920
ttgcttcaac	ctgccgaagc	accgtccgtt	gaggaaaata	taacggaaac	cgttgccgaa	1980
acacccgact	tcaacgccac	cgcagacgat	ttgtccgcat	tacttcaacc	ttctaaagta	2040
cctgccgttg	aggaaaatgc	agcggaaacc	gttgccgatg	atttgtccgc	actgttgcaa	2100
cctgctgaag	caccggccgt	tgaggaaaat	gtaacggaaa	ccgttgccga	aacacccgat	2160
ttcaacgcca	ccgcagacga	tttgtccgca	ttacttcaac	cttctgaagc	acctgccgtt	2220
gaggaaaatg	cagcggaaac	cgttgccgat	gatttgtccg	cactgttgca	acctgctgaa	2280
gcaccggccg	ttgaggaaaa	tgcagcggaa	atcactttgg	aaacgcctga	ttccaacacc	2340
tctgaggcag	acgctttgcc	cgacttcctg	aaagacggcg	aggaggaaac	ggtagattgg	2400
agcatctacc	tctcggaaga	aaatatccca	aataatgcag	ataccagttt	cccttcggaa	2460
tctgtaggtt	ctgacgcgcc	ttccgaagcg	aaatacgacc	ttgccgaaat	gtatctcgaa	2520
atcggcgacc	gcgatgccgc	tgccgagaca	gtgcagaaat	tgctggaaga	agcggaaggc	2580

<210> 2

<211> 875

<212> PRT

<213> Neisseria meningitidis (group B)

<400> 2

Met Pro Ala Gly Arg Leu Pro Arg Arg Cys Pro Met Met Thr Lys Phe 1 5 10 15

Thr Asp Cys Thr Arg Ser Asn Arg Ile Gln Pro Pro Thr His Arg Gly
20 25 30

Tyr Ile Leu Lys Asn Asn Arg Gln Ile Lys Leu Ile Ala Ala Ser Val\$35\$ 40 45

Ala Val Ala Ala Ser Phe Gln Ala His Ala Gly Leu Gly Gly Leu Asn 50 55 60

Ile Gln Ser Asn Leu Asp Glu Pro Phe Ser Gly Ser Ile Thr Val Thr 65 70 75 80

Gly Glu Glu Ala Lys Ala Leu Leu Gly Gly Gly Ser Val Thr Val Ser

85

90

95

Glu Lys Gly Leu Thr Ala Lys Val His Lys Leu Gly Asp Lys Ala Val 100 105 110

Ile Ala Val Ser Ser Glu Gln Ala Val Arg Asp Pro Val Leu Val Phe 115 120 125

Arg Ile Gly Ala Gly Ala Gln Val Arg Glu Tyr Thr Ala Ile Leu Asp 130 135 140

Pro Val Gly Tyr Ser Pro Lys Thr Lys Ser Ala Leu Ser Asp Gly Lys
145 150 155 160

Thr His Arg Lys Thr Ala Pro Thr Ala Glu Ser Gln Glu Asn Gln Asn 165 170 175

Ala Lys Ala Leu Arg Lys Thr Asp Lys Lys Asp Ser Ala Asn Ala Ala 180 185 190

Val	Lys	Pro 195	Ala	Tyr	Asn	Gly	Lys 200	Thr	His	Thr	Val	Arg 205	Lys	Gly	Glu
Thr	Val 210	Lys	Gln	Ile	Ala	Ala 215	Ala	Ile	Arg	Pro	Lys 220	His	Leu	Thr	Leu
Glu 225	Gln	Val	Ala	Asp	Ala 230	Leu	Leu	Lys	Ala	Asn 235	Pro	Asn	Val	Ser	Ala 240
His	Gly	Arg	Leu	Arg 245	Ala	Gly	Ser	Val	Leu 250	His	Ile	Pro	Asn	Leu 255	Asn
Arg	Ile	Lys	Ala 260	Glu	Gln	Pro	Lys	Pro 265	Gln	Thr	Ala	Lys	Pro 270	Lys	Ala
Glu	Thr	Ala 275	Ser	Met	Pro	Ser	Glu 280	Pro	Ser	Lys	Gln	Ala 285	Thr	Val	Glu
Lys	Pro 290	Val	Glu	Lys	Pro	Glu 295	Ala	Lys	Val	Ala	Ala 300	Pro	Glu	Ala	Lys
Ala 305	Glu	Lys	Pro	Ala	Val 310	Arg	Pro	Glu	Pro	Val 315	Pro	Ala	Ala	Asn	Thr 320
Ala	Ala	Ser	Glu	Thr 325	Ala	Ala	Glu	Ser	Ala 330	Pro	Gln	Glu	Ala	Ala 335	Ala
Ser	Ala	Ile	Asp 340	Thr	Pro	Thr	Asp	Glu 345	Thr	Gly	Asn	Ala	Val 350	Ser	Glu
Pro	Val	Glu 355	Gln	Val	Ser	Ala	Glu 360	Glu	Glu	Thr	Glu	Ser 365	Gly	Leu	Phe
Asp	Gly 370	Leu	Phe	Gly	Gly	Ser 375	Tyr	Thr	Leu	Leu	Leu 380	Ala	Gly	Gly	Gly
Ala 385	Ala	Leu	Ile	Ala	Leu 390	Leu	Leu	Leu	Leu	Arg 395	Leu	Ala	Gln	Ser	Lys 400
Arg	Ala	Arg	Arg	Thr	Glu	Glu	Ser	Val	Pro	Glu	Glu	Glu	Pro	Asp	Leu

Asp	Asp	Ala	Ala 420	Asp	Asp	Gly	Ile	Glu 425	Ile	Thr	Phe	Ala	Glu 430	Val	Glu
Thr	Pro	Ala 435	Thr	Pro	Glu	Pro	Ala 440	Pro	Lys	Asn	Asp	Val 445	Asn	Asp	Thr
Leu	Ala 450	Leu	Asp	Gly	Glu	Ser 455	Glu	Glu	Glu	Leu	Ser 460	Ala	Lys	Gln	Thr
Phe 465	Asp	Val	Glu	Thr	Asp 470	Thr	Pro	Ser	Asn	Arg 475	Ile	Asp	Leu	Asp	Phe 480
Asp	Ser	Leu	Ala	Ala 485	Ala	Gln	Asn	Gly	Ile 490	Leu	Ser	Gly	Ala	Leu 495	Thr
Gln	Asp	Glu	Glu 500	Thr	Gln	Lys	Arg	Ala 505	Asp	Ala	Asp	Trp	Asn 510	Ala	Ile
Glu	Ser	Thr 515	Asp	Ser	Val	Tyr	Glu 520	Pro	Glu	Thr	Phe	Asn 525	Pro	Tyr	Asn
Pro	Val 530	Glu	Ile	Val	Ile	Asp 535	Thr	Pro	Glu	Pro	Glu 540	Ser	Val	Ala	Gln
Thr 545	Ala	Glu	Asn	Lys	Pro 550	Glu	Thr	Val	Asp	Thr 555	Asp	Phe	Ser	Asp	Asn 560
Leu	Pro	Ser	Asn	Asn 565	His	Ile	Gly	Thr	Glu 570	Glu	Thr	Ala	Ser	Ala 575	Lys
Pro	Ala	Ser	Pro 580	Ser	Gly	Leu	Ala	Gly 585	Phe	Leu	Lys	Ala	Ser 590	Ser	Pro
Glu	Thr	Ile 595	Leu	Glu	Lys	Thr	Val 600	Ala	Glu	Val	Gln	Thr 605	Pro	Glu	Glu
Leu	His 610	Asp	Phe	Leu	Lys	Val 615	Tyr	Glu	Thr	Asp	Ala 620	Val	Ala	Glu	Thr
Ala 625	Pro	Glu	Thr	Pro	Asp 630	Phe	Asn	Ala	Ala	Ala 635	Asp	Asp	Leu	Ser	Ala 640

Leu	Leu	Gln	Pro	Ala 645	Glu	Ala	Pro	Ser	Val 650	Glu	Glu	Asn	Ile	Thr 655	Glu
Thr	Val	Ala	Glu 660	Thr	Pro	Asp	Phe	Asn 665	Ala	Thr	Ala	Asp	Asp 670	Leu	Ser
Ala	Leu	Leu 675	Gln	Pro	Ser	Lys	Val 680	Pro	Ala	Val	Glu	Glu 685	Asn	Ala	Ala
Glu	Thr 690	Val	Ala	Asp	Asp	Leu 695	Ser	Ala	Leu	Leu	Gln 700	Pro	Ala	Glu	Ala
705					710					715				Pro	720
				725					730					Ser 735	
			740					745					750	Asp	
		755					760					765		Ala	
	770					775					780			Asp	
785 Ser	Ile	Tyr	Leu	Ser	790 Glu	Glu	Asn	Ile	Pro	795 Asn	Asn	Ala	Asp	Thr	800 Ser
Phe	Pro	Ser	Glu	805 Ser	Val	Gly	Ser	Asp	810 Ala	Pro	Ser	Glu	Ala	815 Lys	Tyr
Asp	Leu	Ala	820 Glu	Met	Tyr	Leu	Glu	825 Ile	Gly	Asp	Arg	Asp	830 Ala	Ala	Ala
		835					840		_			845		Leu	
Jiu	850	·	C ±11	, _	Leu	855	J.Lu	Jiu		Jiu	860		·	Leu	_, 5

865 870 875

<210> 3 <211> 483 <212> DNA <213> Neisseria meningitidis (group B) <400> 3 atggaaagga acggtgtatt tggtaaaatt gtcggcaatc gcatactccg tatgtcgtcc gaacacgctg ccgcatccta tccgaaaccg tgcaaatcgt ttaaactagc gcaatcttgg ttcagagtgc gaagctgtct gggcggcgtt tttatttacg gagcaaacat gaaacttatc tataccgtca tcaaaatcat tatcctgctg ctcttcctgc tgcttgccgt cattaatacg gatgccgtta ccttttccta cctgccgggg caaaaattcg atttgccgct gattgtcgta ttgttcggcg catttgtagt cggtattatt tttggaatgt ttgccttgtt cggacggttg ttgtcgttac gtggcgagaa cggcaggttg cgtgccgaag taaagaaaaa tgcgcgtttg acggggaagg agctgaccgc accaccggcg caaaatgcgc ccgaatctac caaacagcct taa <210> 4 <211> 160 <212> PRT <213> Neisseria meningitidis (group B) <400> 4 Met Glu Arg Asn Gly Val Phe Gly Lys Ile Val Gly Asn Arg Ile Leu 5 1.0 Arg Met Ser Ser Glu His Ala Ala Ser Tyr Pro Lys Pro Cys Lys 20 25 Ser Phe Lys Leu Ala Gln Ser Trp Phe Arg Val Arg Ser Cys Leu Gly 35 40 Gly Val Phe Ile Tyr Gly Ala Asn Met Lys Leu Ile Tyr Thr Val Ile 50 55 60 Lys Ile Ile Ile Leu Leu Phe Leu Leu Leu Ala Val Ile Asn Thr 65 70 75

60

120

180

240

300

360

420

480

483

Asp Ala Val Thr Phe Ser Tyr Leu Pro Gly Gln Lys Phe Asp Leu Pro 85 90 95

Leu Ile Val Val Leu Phe Gly Ala Phe Val Val Gly Ile Ile Phe Gly 105

Met Phe Ala Leu Phe Gly Arg Leu Leu Ser Leu Arg Gly Glu Asn Gly 115 120

Arg Leu Arg Ala Glu Val Lys Lys Asn Ala Arg Leu Thr Gly Lys Glu 135 130 140

Leu Thr Ala Pro Pro Ala Gln Asn Ala Pro Glu Ser Thr Lys Gln Pro 145 150 155 160

<210> 5

<211> 1551

<212> DNA

<213> Neisseria meningitidis (group B)

<400> 5

atgaaaaaac ctttgatttc ggttgcggca gcattgctcg gcgttgcttt gggcacgcct tattatttgg gtgtcaaagc cgaagaaagc ttgacgcagc agcaaaaaat attgcaggaa 120 180 acgggcttct tgaccgtcga atcgcaccaa tatgagcgcg gctggtttac ctctatggaa acgacggtca tccgtctgaa acccgagttg ctgaataatg cccgaaaata cctgccggat 240 aacctgaaaa cagtgttgga acagccggtt acgctggtta accatatcac gcacggccct 300 ttcgccggcg gattcggcac gcaggcgtac attgaaaccg agttcaaata cgcgcctgaa 360 acggaaaaag ttctggaacg cttttttgga aaacaagtcc cggcttccct tgccaatacc 420 gtttatttta acggcagcgg taaaatggaa gtcagtgttc ccgccttcga ttatgaagag 480 540 ctgtcgggca tcaggctgca ctgggaaggc ctgacgggag aaacggttta tcaaaaaggt ttcaaaagct accggaacgg ctatgatgcc cccttgttta aaatcaagct ggcagacaaa 600 ggcgatgccg cgtttgaaaa agtgcatttc gattcggaaa cttcagacgg catcaatccg 660 cttgctttgg gcagcagcaa tctgaccttg gaaaaattct ccctagaatg gaaagagggt 720 gtcgattaca acgtcaagtt aaacgaactg gtcaatcttg ttaccgattt gcagattggc 780 gcgtttatca atcccaacgg cagcatcgca ccttccaaaa tcgaagtcgg caaactggct 840 ttttcaacca agaccgggga atcaggcgcg tttatcaaca gtgaagggca gttccgtttc 900 gatacactgg tgtacggcga tgaaaaatac ggcccgctgg acatccatat cgctgccgaa 960 1020

cacctcgatg cttctgcctt aaccgtattg aaacgcaagt ttgcacaaat ttccgccaaa

aaaatgaccg	aggaacaaat	ccgcaatgat	ttgattgccg	ccgtcaaagg	agaggcttcc	1080
ggactgttca	ccaacaatcc	cgtattggac	attaaaactt	tccgattcac	gctgccatcg	1140
ggaaaaatcg	atgtgggcgg	aaaaatcatg	tttaaagaca	tgaagaagga	agatttgaat	1200
caattgggtt	tgatgctgaa	gaaaaccgaa	gccgacatca	gaatgagtat	tccccaaaaa	1260
atgctggaag	acttggcggt	cagtcaagca	ggcaatattt	tcagcgtcaa	tgccgaagat	1320
gaggcggaag	gcagggcaag	tcttgacgac	atcaacgaga	ccttgcgcct	gatggtggac	1380
agtacggttc	agagtatggc	aagggaaaaa	tatctgactt	tgaacggcga	ccagattgat	1440
actgccattt	ctctgaaaaa	caatcagttg	aaattgaacg	gtaaaacgtt	gcaaaacgaa	1500
ccggagccgg	attttgatga	aggcggtatg	gtttcagagc	cgcagcagta	a	1551

<210> 6

<211> 516

<212> PRT

<213> Neisseria meningitidis (group B)

<400> 6

Met Lys Lys Pro Leu Ile Ser Val Ala Ala Ala Leu Leu Gly Val Ala 1 5 10 15

Leu Gly Thr Pro Tyr Tyr Leu Gly Val Lys Ala Glu Glu Ser Leu Thr 20 25 30

Gln Gln Gln Lys Ile Leu Gln Glu Thr Gly Phe Leu Thr Val Glu Ser 35 40 45

His Gln Tyr Glu Arg Gly Trp Phe Thr Ser Met Glu Thr Thr Val Ile 50 60

Arg Leu Lys Pro Glu Leu Leu Asn Asn Ala Arg Lys Tyr Leu Pro Asp 65 70 75 80

Asn Leu Lys Thr Val Leu Glu Gln Pro Val Thr Leu Val Asn His Ile \$85\$ 90 95

Thr His Gly Pro Phe Ala Gly Gly Phe Gly Thr Gln Ala Tyr Ile Glu 100 105 110

Thr Glu Phe Lys Tyr Ala Pro Glu Thr Glu Lys Val Leu Glu Arg Phe 115 120 125

Phe Gly Lys Gln Val Pro Ala Ser Leu Ala Asn Thr Val Tyr Phe Asn 135 Gly Ser Gly Lys Met Glu Val Ser Val Pro Ala Phe Asp Tyr Glu Glu 145 150 155 160 Leu Ser Gly Ile Arg Leu His Trp Glu Gly Leu Thr Gly Glu Thr Val 165 170 175 Tyr Gln Lys Gly Phe Lys Ser Tyr Arg Asn Gly Tyr Asp Ala Pro Leu 180 185 190 Phe Lys Ile Lys Leu Ala Asp Lys Gly Asp Ala Ala Phe Glu Lys Val 195 200 205 His Phe Asp Ser Glu Thr Ser Asp Gly Ile Asn Pro Leu Ala Leu Gly 210 220 215 Ser Ser Asn Leu Thr Leu Glu Lys Phe Ser Leu Glu Trp Lys Glu Gly 225 230 235 240

Val Asp Tyr Asn Val Lys Leu